

STIC Search Report Biotech-Chem Library

STIC Deletes Trend no vi

TO: Minh-Tam Davis

Art Unit: 1642

Location: rem/3A24/3C18 Serial Number: 10/057813

Friday, October 07, 2005

From: Beverly Shears

Location: Biotech-Chem Library

REM 1A54

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Sexion Notes

Protein Sequence Searches – February 2005

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (uniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.



STIC-Biotech/ChemLib

167306

From:

PTO-1590 (9-90)

Davis, Minh-Tam

Sent:

Thursday, September 29, 2005 1:46 PM STIC-Biotech/ChemLib

To: Subject:

search request for 10/057813

Please search in commercial database, issued patent files, PGPUB and interference:

1) Amino acids 1-91, and 85-125 of SEQ ID NO:14, with size limitation for the sequences in the database to the size of the amino acid fragments.

Thank you.
MINH TAM DAVIS
ART UNIT 1642, ROOM 3A24, MB 3C18
272-0830

seg 14-163AA

SEP 29 2005

ALVOHENTERS

Searcher: Searcher Phone: Date Searcher Picked up: Date completed: Searcher Prep Time: Online Time:	Type of Search NA# AA#: S/L: Oligomer: Encode/Transl: Structure #:Text: Inventor: Litigation:	Vendors and cost where applicable STN: DIALOG: QUESTEL/ORBIT: LEXIS/NEXIS: SEQUENCE SYSTEM:
		WWW/Internet:Other (Specify):
Date completed:	Search Site	Vendors
Searcher: Bevering e	STIC	IG
Terminal time:	CM-1	STN
Elapsed time:	•	Dialog
CPU time:	Type of Search	APS .
. Total time:	N.A. Sequence	Geninfo
Number of Searches:	A.A. Sequence	SDC
Number of Databases:	Structure :	DARC/Questel
	Bibliographic	Other